

Angiotensin II stimulates the synthesis of vascular endothelial growth factor through the p38 mitogen activated protein kinase pathway in cultured mouse podocytes

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Abstract

Angiotensin II (Ang-II) and vascular endothelial growth factor (VEGF) have an important role in the pathogenesis of diabetic nephropathy, but the signaling cascade of VEGF regulation in response to Ang-II in podocytes is largely unknown. In these experiments, we looked at the effect of Ang-II on the production of VEGF, and investigated whether VEGF production depends on the p38 mitogen activated protein kinase (MAPK) pathway in cultured mouse podocytes. Incubation of podocytes with Ang-II induced a rapid increase in VEGF mRNA expression and protein synthesis as well as its transcriptional activity in an Ang-II dose-dependent manner. To further define the role of angiotensin type 1 (AT1) and type 2 (AT2) receptors involved in Ang-II-mediated VEGF synthesis, the effects of selective AT1 and AT2 receptor antagonists were evaluated. Prior treatment with losartan significantly inhibited VEGF mRNA and protein synthesis induced by Ang-II, which suggests that the AT1 receptor is involved in Ang-II-mediated VEGF synthesis. Furthermore, stimulation of the cells with Ang-II increased both phosphorylation of p38 MAPK and MAP kinase kinase 3/6 (MKK3/6). Additionally, Ang-II enhanced the DNA binding activity to cAMP response element binding protein (CREB) and phosphorylation of CREB. In addition, to investigate the role of p38 MAPK in Ang-II-induced VEGF synthesis, podocytes were pretreated with or without the p38 MAPK inhibitor, SB203580 for 24 h to observe whether Ang-II-mediated VEGF synthesis was inhibited by blocking p38 MAPK. The addition of SB203580 led to a marked inhibition of the increased VEGF mRNA and protein production induced by Ang-II in a dose-dependent manner. Taken together, these results suggest that Ang-II stimulates the synthesis of VEGF in podocytes and the production of VEGF induced by Ang-II is mediated, in part, through the activation of the p38 MAPK pathway.

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Introduction

Diabetic milieu results in the increased expression of angiogenic growth factors in numerous tissues as a response to both hyperglycemia and tissue ischemia (Duh & Aiello 1999, Ferrara 1999). It is generally accepted that podocytes are the major site of vascular endothelial growth factor (VEGF) production, and their regulation in VEGF expression may play an important role in diabetic nephropathy (Brown *et al.* 1992, Kretzler *et al.* 1998). Podocytes are unique cells in the glomerulus, cover the outer part of glomerular basement membrane and function as the final filtration barrier to protein loss.

There is now increasing data supporting a direct role for VEGF in the pathogenesis of diabetic nephropathy. VEGF is up-regulated from the early stage of diabetic

nephropathy (Cooper *et al.* 1999). *In vivo* blockade of VEGF abolished hyperfiltration and suppressed the urinary albumin excretion rate in diabetic rats (De Zeeuw *et al.* 2001). In addition, VEGF may contribute to mesangial expansion and phagocytes infiltration (Clausz *et al.* 1990, Flyvbjerg *et al.* 2002). Taken together, these studies suggest that VEGF plays an important role in the pathogenesis of diabetic nephropathy.

Angiotensin II (Ang-II) plays an important role in the development of glomerulosclerosis (Leehey *et al.* 2000). There are several reports showing that Ang-II increases VEGF production in human mesangial cells and in vascular smooth muscle cells (Williams *et al.* 1995, Gruden *et al.* 1999, Pupilli *et al.* 1999). However, there is no report on whether Ang-II induces VEGF production in podocytes, which are the major source of VEGF

synthesis in the glomerulus. Additionally, the signaling cascade of VEGF regulation in response to Ang-II in podocytes is largely unknown. Many studies have demonstrated protein kinase C activation in diabetic glomeruli and mesangial cells cultured under high glucose conditions (Craven & DeRubertis 1989, Williams & Schrier 1993, Ziyadeh *et al.* 1995, Hoshi *et al.* 2002). Increased p38 mitogen activated protein kinase (MAPK) activity has been demonstrated in diabetic glomeruli (Kang *et al.* 2001), and Ang-II also activated p38 MAPK leading to the induction of fibronectin expression and cellular growth in rat mesangial cells (Reddy *et al.* 2002).

In this study, we examined the effects of Ang-II on VEGF mRNA expression and protein production in cultured mouse podocytes, and evaluated whether VEGF production depends on the p38 MAPK pathway. In addition, we examined the relationship between the p38 MAPK pathway and VEGF synthesis in cultured mouse podocytes.

Materials and methods

Cell cultures

A thermosensitive, SV40-transfected immortalized mouse podocyte cell line, a generous gift from Peter Mundel (Albert Einstein College of Medicine, New York, NY, USA), was used for this study. Cultivation of mouse podocytes that were conditionally immortalized with a temperature-sensitive variant of the SV40 large T antigen (tsA58) and whose activity can be increased by γ -interferon, was performed as described previously (Mundel *et al.* 1997). To propagate podocytes, cells were cultivated at 33 °C and treated with 10–50 U/ml mouse recombinant γ -interferon (permissive condition), which increases the expression of the temperature-sensitive large T antigen and cell proliferation. To induce differentiation, podocytes were thermoshifted to 37 °C and deprived of γ -interferon (non-permissive condition) for 14 days. Studies were performed using a podocytes cell line at 20–24 passages. Identification of podocytes was performed using RT-PCR for podocyte specific markers such as Wilms' tumor protein (WT-1). Differentiation of podocytes was determined on the basis of the expression of synaptopodin, which is a differentiation marker, using RT-PCR.

Experimental design

Differentiated podocytes were grown to sub-confluence in type 1 collagen-coated dishes (Iwaki, Tokyo, Japan) in growth media containing 10% fetal calf serum (FCS), 100 U/ml penicillin, and 0.1 mg/ml streptomycin in RPMI 1640 medium, and then cultured for 24 h in a medium containing 5 mmol/l D-glucose and 1% FCS

before being exposed to experimental conditions. In the Ang-II-stimulated group, different concentrations of Ang-II were added to the culture media at final concentrations of 1 nM, 10 nM, and 100 nM. To define the mechanism of Ang-II-mediated VEGF synthesis, we used both the Ang-II receptor type 1 antagonist (losartan) and the Ang-II receptor type 2 antagonist (PD123319). When the effects of these drugs were tested, these compounds were added to podocytes one hour before Ang-II treatment. All experimental groups were cultured in triplicate and harvested at 6, 24 and 72 h for extraction of total RNA and protein. The results are representative of those from three independent experiments.

To determine the effect of Ang-II on the p38 MAPK pathway, sub-confluent podocytes were cultured for 24 h in medium containing 5 mmol/l D-glucose and 1% FCS. Then, the cells were treated with Ang-II at a final concentration of 100 nM, and harvested after 30 min, 1 h, 6 h and 24 h. In studies examining the role of the p38 MAPK pathway on VEGF synthesis, SB203580, which is a p38 MAPK inhibitor, was added to podocytes one hour before Ang-II treatment at final concentrations of 1 and 10 μ M. Cells were harvested at 30 min, 1 h, 6 h, and 24 h and then total RNA and protein were extracted. To avoid any confounding effects of serum on VEGF and proteins in the p38 MAPK pathway experiments, serum-free media were used. All experimental groups were cultured in triplicate. The results are representative of those from three independent experiments.

RT-PCR

Total RNA extraction was performed with a Trizol reagent and the cDNA was synthesized by a reverse transcription reaction using an RNA PCR kit (Applied Biosystems, Roche Inc., Foster City, CA, USA) in a 20 μ l mixture containing 1 μ g RNA, 50 mM KCl, 10 mM Tris/HCl, 5 mM MgCl₂, 1 mM of each dNTPs, oligo-(dT) primers, 20 units RNase inhibitor and 50 units Moloney Murine Leukemia Virus (MuLV) reverse transcriptase. The reaction mixture was incubated for 60 min at 42 °C, then heated at 90 °C for 7 min in a thermocycler (GeneAmp PCR system 9600, Perkin Elmer, Roche Molecular System, Branchburg, NJ, USA). Next, cDNA was amplified by 2.5 units AmpliTaq Gold polymerase in a 25 μ l reaction volume containing 10 mmol/l Tris-HCl (pH 8.3), 50 mmol/l KCl, 1.5 mmol/l MgCl₂, 0.2 mmol/l deoxynucleoside triphosphate, and 30 pmol of each primer. Sequence specific primers for VEGF, designed according to the mouse sequence (Gene bank number M95200), were used for the PCR. The nucleotide sequences of each primer are as follows: sense 5'-CAGGCTGCTGT AACGATGAA-3' and antisense 5'-AATGCTTCTC CGCTCTGAA-3'. As an internal control, β -actin was also amplified and nucleotide sequences for primers are

as follows: sense 5'-TCATGAGGTAGTCCGTCAGG-3' and antisense 5'-TCTAGGCACCAAGGTGTG-3'. The PCR conditions were 40 cycles of denaturation at 95 °C for 45 s, annealing at 58 °C for 45 s, and extension at 72 °C for 3 min. The number of PCR cycles selected represents a point before the plateau of amplification products, as described previously (Cha *et al.* 2000, Kim *et al.* 2000). To confirm the identity of each PCR product, each of the electrophoresed PCR bands were extracted with a DNA extraction kit (Qiagen, Valencia, CA, USA) and sequenced using an ABI automated DNA sequencing system (ABI Genetic Analyzer 310; PRISM, Branchburg Park, NJ, USA). The RT-PCR products were separated on a 2% agarose gel with ethidium bromide staining by electrophoresis. After scanning at 300 d.p.i., densitometric analysis was performed for quantification using NIH (Bethesda, MA, USA) image analysis software (version 1.61). The ratios of the concentration of β -actin to those of VEGF were evaluated.

Transient transfection and luciferase reporter activity assay

VEGF reporter plasmid containing firefly luciferase linked to VEGF promoter sequences was a generous gift from Dr P A D'Amore (Shima *et al.* 1996). Cells were plated onto 24-well plates at a density of 1×10^5 cells/well. Following 24 h growth, when cells were approximately 40 to 50% confluence, cells were transfected with 1 μ g VEGF reporter plasmid and 1 μ g plasmid containing Renilla luciferase driven by TK promoter using Superfect, as recommended by the manufacturer (Qiagen) for 24 h. Then, media containing the transfection reagent were replaced with complete media without serum, and cells were treated with or without different concentrations of angiotensin II at final concentrations of 1 nM, 10 nM and 100 nM. After 24 h, luciferase activity was determined using the dual luciferase assay system according to the manufacturer's instructions (Promega Corp., Madison, WI, USA). Relative light units from firefly luciferase activity were measured using a luminometer (Mono Light 2010, Analytical Luminescence Lab., San Diego, CA, USA). To control for differences in transfection efficiency from well to well, plasmid containing Renilla luciferase driven by TK promoter was included in each transfection and used for normalization.

Western blotting

Cells from the three experiments were lysed in lysis buffer (150 mM NaCl, 50 mM Tris-HCl, pH 8.0, 1 mM phenylmethylsulfonylfluoride and 1% Triton X-100) with a sonicator, and total protein concentration was measured by the method of Bradford (Bio-Rad). Thirty micrograms protein were electrophoresed on a 10%

sodium dodecyl sulfate-polyacrylamide gel under denaturing conditions. The proteins were transferred onto a polyvinylidene difluoride (PVDF) membrane (Immobilon-P, Millipore, Bedford, MA, USA) for 150 min at 250 mA. After the filter was blocked by incubating the membranes with blocking solution ($1 \times$ PBS, 0.15% Tween 20 and 5% non-fat milk) for 2 h at room temperature, the membranes were hybridized with each specific primary antibody overnight at 4 °C. Anti-rabbit polyclonal VEGF antibody (Santa Cruz Biotechnology, Santa Cruz, CA, USA), polyclonal antibody to p38 MAPK, phospho-specific p38 MAPK, MAP kinase kinase 3/6 (MKK3/6), phospho-specific MKK3/6, cAMP response element binding protein (CREB), and phospho-specific CREB (New England Biolabs, Inc., Beverly, MA, USA) diluted 1:1000 were applied to the membrane. The filter was then washed four times with phosphate buffered saline Tween-20 (PBST), and incubated with horseradish peroxidase-conjugated secondary antibody diluted 1:1000 for 60 min at room temperature. The detection of specific signals was performed using the ECL method (Amersham Biosciences Corp., Piscataway, NJ, USA). Equal amounts of protein loading were confirmed by Coomassie Blue staining of the gel.

Enzyme-linked immunosorbent assay (ELISA)

The amount of VEGF protein secreted by podocytes in culture medium was measured by commercially available quantitative sandwich enzyme immunoassay (R&D Systems, Minneapolis, MN, USA). The assay method was designed to recognize both the 164 and 120 amino acid residue forms of mouse VEGF. The sensitivity of ELISA for VEGF was 3 pg/ml. The intra-assay coefficient of variation was 4.7% and the interassay coefficient of variation was 6.4%. For measurement of secreted soluble VEGF in cultured medium, conditioned media were collected at the end of the treatment periods. All particulates were then removed by centrifugation at $4000 \times g$ for 10 min and stored at -20 °C before the measurement of VEGF proteins. Supernatants were diluted five-fold with diluent solution according to the manufacturer's instructions. Fifty microliters of sample with an equal volume of diluent were dispensed in a 96-microwell plate, precoated with polyclonal antibody specific for mouse VEGF. The plates were incubated at room temperature for 2 h, washed 5 times, and developed with 100 μ l color reagent per well. The intensity of the color was measured in an ELISA reader at 540 nm.

Electrophoretic mobility shift assay (EMSA)

Podocytes were treated with 100 nM angiotensin II for 8 h. Nuclear extracts were prepared by the method of Dignam *et al.* (1983). Briefly, nuclear extracts (5 μ g

protein) were preincubated with poly(dI-dC)zpoly(dI-dC) (2 µg), dithiothreitol (0.3 mM), and reaction buffer (12 mM Tris, pH 7.9, 2 mM MgCl₂, 60 mM KCl, 0.12 mM EDTA, and 12% glycerol) with or without CREB antiserum (1–2 µl) for 30 min at 4 °C. ³²P-Labeled oligonucleotide containing a consensus cAMP response element (CRE) sequence (5'-AGAGATTGCC TGACGTCAGAGAGCTAG-3', Promega) was then added, and the reaction mixtures were incubated for 10 min at 37 °C. The reaction mixtures were then separated on a 4% nondenaturing polyacrylamide gel at 200 V for 2 h. The gel was dried and autoradiographed. Anti-CREB antibody (Santa Cruz Biotechnology) was used for supershift. Unlabeled double-stranded oligonucleotides with one copy of CRE or Oct1 (5'-TGTCGAA

TGCAAATCACTAGAA-3', Promega) were used as competitor.

Statistical analysis

We used non-parametric analysis due to the small sample number. Results are expressed as means ± s.d. Kruskal-Wallis was used for comparison of more than two groups, followed by Mann-Whitney *U* test for comparison, using a microcomputer-assisted program with SPSS for Windows 10.0 (SPSS Inc., Chicago, IL, USA). *P* < 0.05 was considered statistically significant.

Results

Effects of Ang-II on VEGF mRNA and protein production

When different concentrations of Ang-II were added to podocytes for 6 h, VEGF mRNA expression incrementally increased in an Ang-II concentration-dependent manner compared with controls (1 nM 0.43 ± 0.01; 10 nM 0.95 ± 0.10; 100 nM 1.95 ± 0.07; control 0.28 ± 0.04). For Ang-II concentrations of 1, 10 and 100 nM, VEGF mRNA levels were time-dependently increased at 6 h and 24 h compared with that of the control group (group with Ang-II at 100 nM: 6 h 1.95 ± 0.07, 24 h 4.25 ± 0.62, 72 h 1.32 ± 0.09). As shown in Fig. 1, the stimulatory effect of Ang-II on VEGF mRNA levels showed significant increases at 6 h and 24 h, and decreased at 72 h, although its level was still higher compared with the control group.

To define the role of angiotensin type 1 (AT1) and type 2 (AT2) receptors involved in Ang-II-mediated

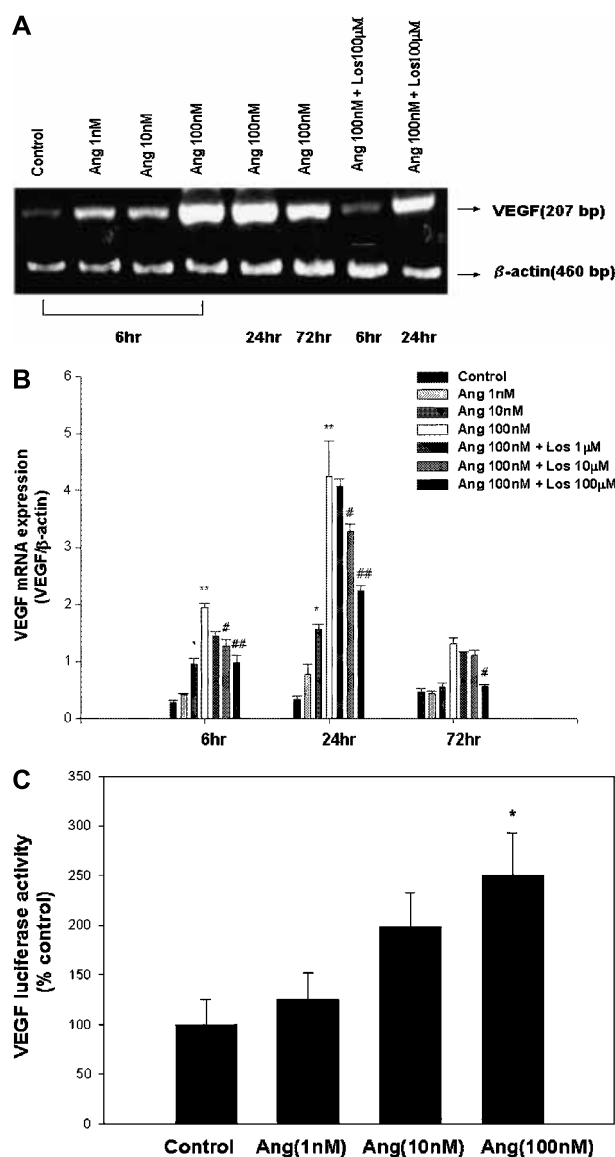


Figure 1 Effects of Ang-II and Ang-II type 1 receptor blockade (losartan) on the expression of VEGF mRNA and the effect of Ang-II on the transcriptional activity of VEGF in cultured podocytes. (A) Representative RT-PCR showing the 207 bp product, which is identical to the 260 to 266 nucleotide of mouse alternatively spliced VEGF₁₆₄. In addition, a second 279 bp product was also detected, but the density of the band was much weaker than the 207 bp band. The 279 bp band had an additional 72 nucleotides corresponding to the VEGF₁₈₈ isoform. In this figure, only the major band of the RT-PCR product is shown. Podocytes were exposed to different concentrations of Ang-II with or without different concentrations of losartan for 6, 24 and 72 h. (B) Densitometric analysis of RT-PCR data. Results are expressed as an optical density ratio of VEGF/β-actin. Data are shown as means ± s.d. of 3 independent experiments with triplicate dishes. (C) Podocytes were co-transfected with VEGF promoter-driven luciferase vector and TK-driven Renilla luciferase plasmid. Then cells were treated with Ang-II at concentrations of 1, 10 or 100 nM for 24 h. VEGF reporter activity was normalized to Renilla luciferase activity. Data are shown as means ± s.d. *n* = 6 in each group. Ang, angiotensin II; Los, losartan. **P* < 0.05, ***P* < 0.001 vs control. #*P* < 0.05, ##*P* < 0.01 vs group treated with 100 nM Ang-II.

VEGF mRNA upregulation, podocytes were preincubated with losartan, a selective AT1 receptor antagonist and PD123319, a selective AT2 receptor antagonist under the influence of 100 nM Ang-II. Losartan at a concentration of 1 μ M did not show a significant effect on VEGF mRNA expression. However, supplementation with 10 and 100 μ M losartan significantly inhibited the VEGF mRNA expression induced by Ang-II (in the group with 100 μ M losartan: 6 h 0.98 ± 0.14 , 24 h 2.24 ± 0.09 , 72 h 0.56 ± 0.04) (Fig. 1A,B).

Effect of Ang-II on transcriptional activity of VEGF in cultured podocytes

Since Ang-II stimulated VEGF mRNA and protein production, we wished to confirm using a luciferase reporter assay whether VEGF production induced by Ang-II actually increased its transcriptional activity. Podocytes transfected with VEGF-luciferase exhibited an increase in luciferase activity after stimulation with Ang-II in a dose-dependent manner. Although Ang-II treatment did not induce a significant increase in transcriptional activity at a concentration of 1 and 10 nM, VEGF transcriptional activity was significantly increased 2.5-fold at a concentration of 100 nM compared with controls (Fig. 1C).

VEGF protein content in podocytes, evaluated by western blot, also increased in a concentration- and time-dependent manner after 6-h treatment with Ang-II (Fig. 2A,B). Densitometric analysis of the VEGF protein demonstrated a 1.9-fold (at 6 h), 2.5-fold (at 24 h) and 2.7-fold (at 72 h) higher level compared with controls in the Ang-II (100 nM)-treated group. Losartan pretreatment (100 μ M) induced a reduction of 16% at 6 h, 44% at 24 h, and 56% at 72 h in VEGF synthesis induced by Ang-II (100 nM).

To confirm the effect of Ang-II on the upregulation of VEGF protein synthesis, the secretory types of VEGF were measured by the ELISA technique. Ang-II stimulation also increased the release of VEGF protein in a dose- and time-dependent manner. The levels of released VEGF after exposure for 6 h to 1 nM, 10 nM and 100 nM Ang-II (172 ± 18 pg/ 10^5 cells/ml, 222 ± 22 pg/ 10^5 cells/ml, and 342 ± 28 pg/ 10^5 cells/ml respectively) were significantly higher than the control cells (94 ± 11 pg/ 10^5 cells/ml) (Fig. 2C). For an Ang-II concentration of 100 nM, the time course of VEGF release showed 342 ± 28 pg/ 10^5 cells/ml at 6 h, 424 ± 20 pg/ 10^5 cells/ml at 24 h, and 426 ± 18 pg/ 10^5 cells/ml at 72 h compared with the level of controls (6 h 94 ± 11 pg/ 10^5 cells/ml, 24 h 101 ± 10 pg/ 10^5 cells/ml, 72 h 116 ± 12 pg/ 10^5 cells/ml).

Supplementation of culture media with losartan even at a concentration of 1 μ M showed significant inhibition of VEGF release. In comparison with the 100 nM Ang-II treatment group the levels of VEGF release in the group

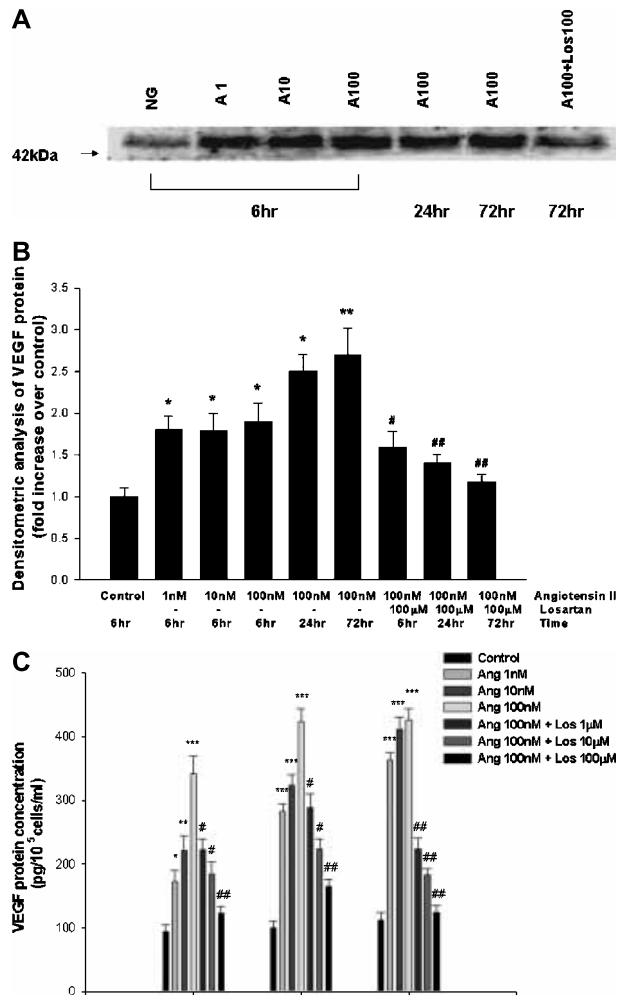


Figure 2 Effect of Ang-II and Ang-II type 1 receptor blockade (losartan) on VEGF protein in cultured podocytes. Podocytes were exposed to different concentrations of Ang-II with or without different concentrations of losartan for 6, 24 and 72 h. (A) VEGF protein in podocytes was detected as a single band of approximately 42 kDa. Thirty micrograms proteins were electrophoresed on 10% sodium dodecyl sulfate-polyacrylamide gel. (B) Densitometric data are shown as means \pm S.D. of 3 independent experiments with triplicate dishes. (C) Secretory VEGF levels from culture supernatants were measured by ELISA. Data are shown as means \pm S.D. of 3 independent experiments with triplicate dishes. Ang, angiotensin II; Los, losartan. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ vs control. # $P < 0.05$, ## $P < 0.01$ vs group treated with 100 nM Ang-II.

with prior treatment with losartan (100 μ M) were significantly inhibited at 6 h (Ang-II 342 ± 28 pg/ 10^5 cells/ml vs Ang-II+losartan 123 ± 11 pg/ 10^5 cells/ml), 24 h (Ang-II 424 ± 20 pg/ 10^5 cells/ml vs Ang-II+losartan 165 ± 11 pg/ 10^5 cells/ml), and 72 h (Ang-II 426 ± 18 pg/ 10^5 cells/ml vs Ang-II+losartan 124 ± 11 pg/ 10^5 cells/ml) (Fig. 2C). However, PD123319,

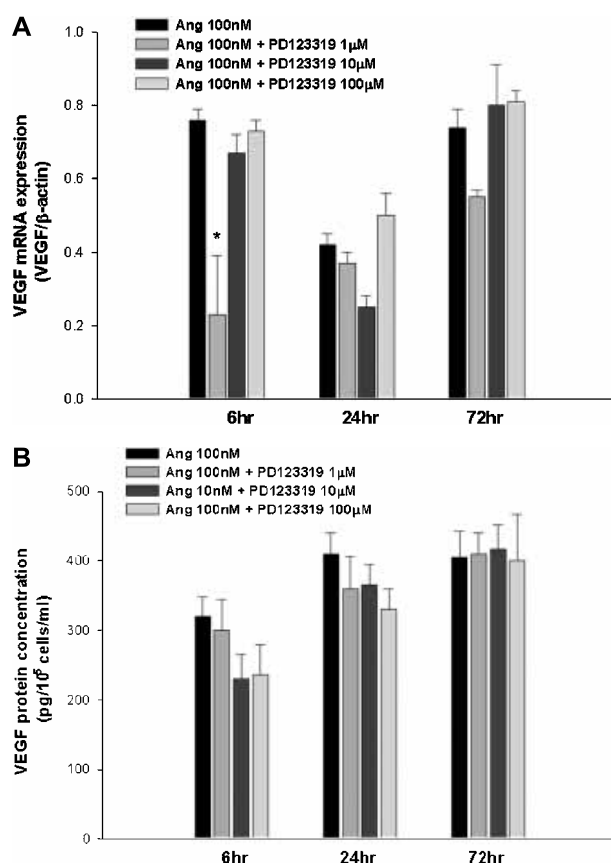


Figure 3 Effects of Ang-II type 2 receptor blockade (PD123319) on the expression of VEGF mRNA and protein secretion in cultured podocytes. Podocytes were exposed to 100 nM Ang-II with or without different concentrations of PD123319 for 6, 24 and 72 h. (A) Densitometric analysis of RT-PCR data. Results are expressed as an optical density ratio of VEGF/β-actin. Data are shown as means±S.D. of 3 independent experiments with triplicate dishes. (B) Secretory VEGF proteins were measured in culture supernatant using ELISA, which recognizes both the 164 and 120 amino acid residue forms of mouse VEGF. Data are shown as means±S.D. of 3 independent experiments with triplicate dishes. Ang, angiotensin II. * $P < 0.05$ vs group treated with 100 nM Ang-II.

which is a selective AT₂ receptor antagonist, did not show any significant effects on Ang-II-induced VEGF mRNA expression and protein production (Fig. 3).

Effects of Ang-II on the activation of p38 MAPK

Since Ang-II stimulated VEGF mRNA and protein production, we further evaluated whether VEGF production induced by Ang-II depends on the p38 MAPK pathway. We first examined the activation of p38 MAPK in response to Ang-II stimulation with 100 nM Ang-II, which was the concentration of Ang-II

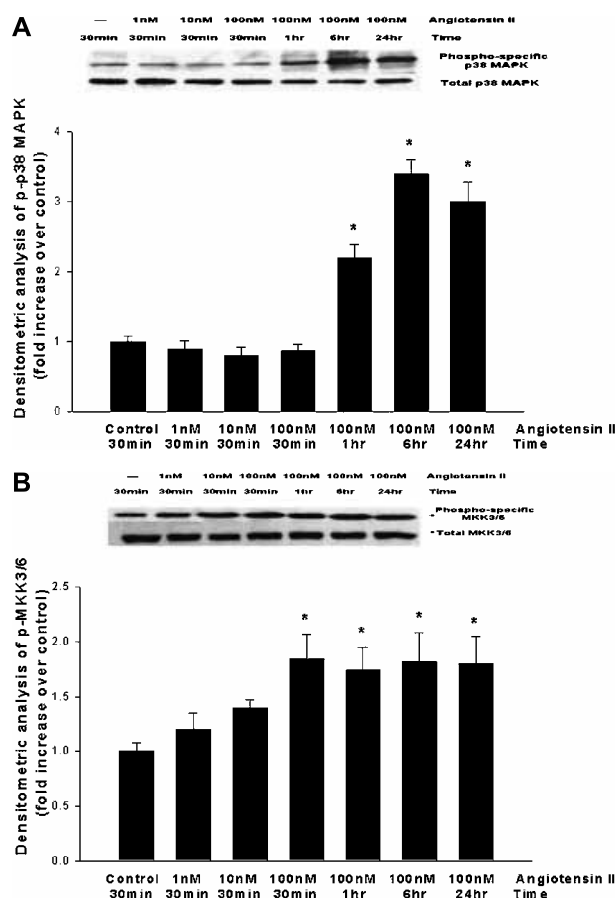


Figure 4 Effect of Ang-II on the activation of p38 MAPK protein and MKK3/6 protein in cultured podocytes. Different concentrations of Ang-II were added to the culture media at final concentrations of 1 nM, 10 nM and 100 nM for 30 min. At 100 nM Ang-II, podocytes were harvested at 30 min, 1 h, 6 h, and 24 h. (A) Representative Western blot of phospho-specific p38 mitogen activated protein kinase (MAPK) protein in cultured podocytes in response to Ang-II treatment at 30 min, 1 h, 6 h and 24 h. (B) Representative Western blot of phospho-specific p38 mitogen activated protein kinase kinase 3/6 (MKK3/6) protein in cultured podocytes in response to Ang-II treatment at 30 min, 1 h, 6 h and 24 h. Densitometric data are shown as means±S.D. of 3 independent experiments with triplicate dishes. * $P < 0.05$ vs control.

that induced maximal production of VEGF synthesis. The activation of p38 MAPK, assessed by measuring the levels of phospho-specific p38 MAPK, was found to increase rapidly in response to Ang-II after one hour. The maximal activity was observed at the 6-h interval, densitometric analysis showed a 3.4 times higher level compared with controls (Fig. 4A). Following this peak, the level of p38 MAPK phosphorylation gradually decreased but remained higher than controls. However, the total p38 MAPK protein level did not differ among the groups.

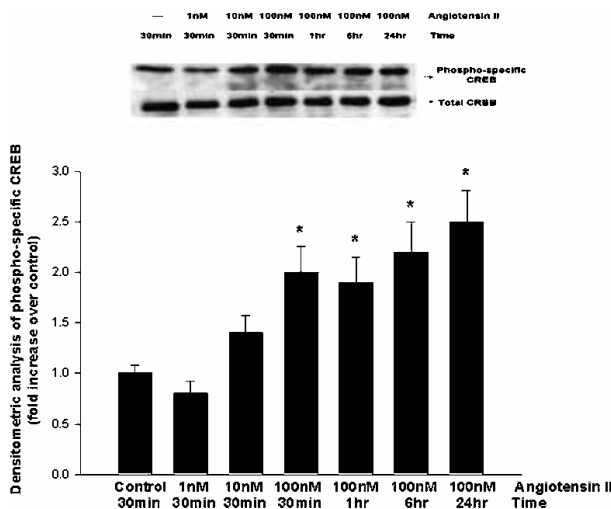


Figure 5 Effect of Ang-II on the activation of CREB protein in cultured podocytes. Different concentrations of Ang-II were added to the culture media at final concentrations of 1 nM, 10 nM and 100 nM for 30 min. At 100 nM Ang-II, podocytes were harvested at 30 min, 1 h, 6 h, and 24 h. Western blot was performed using both total CREB antibody and phospho-specific CREB antibody. Densitometric data are shown as means \pm S.D. of 3 independent experiments with triplicate dishes. * P < 0.05 vs control.

Effects of Ang-II on the activation of MKK3/6

We next examined the effect of Ang-II on the activation of MKK3/6, which is an upstream activator of p38 MAPK. Phospho-specific MKK3/6 levels, an indication of MKK3/6 activation, peaked in response to Ang-II at 30 min and remained at higher levels throughout the study period (1.8-fold on average) (Fig. 4B). The increase in phospho-specific MKK3/6 levels occurred earlier than that in phospho-specific p38 MAPK levels. However, there was no significant difference in total MKK3/6 protein expression among the groups.

Effects of Ang-II on the activation of CREB

To determine whether the activation of the p38 MAPK pathway could induce a parallel increase in the activity of a p38 MAPK target transcription factor, we observed the activation of CREB, which is a known transcription factor for the synthesis of VEGF (Shima *et al.* 1996). CREB activation measured by the level of phospho-specific CREB demonstrated dose- and time-dependent increments after Ang-II stimulation. Maximum activity was observed at 24 h incubation, and densitometric analysis showed 2.5 times higher levels compared with controls (Fig. 5). However, total CREB protein expression was not changed according to the concentration of Ang-II and the stimulation time.

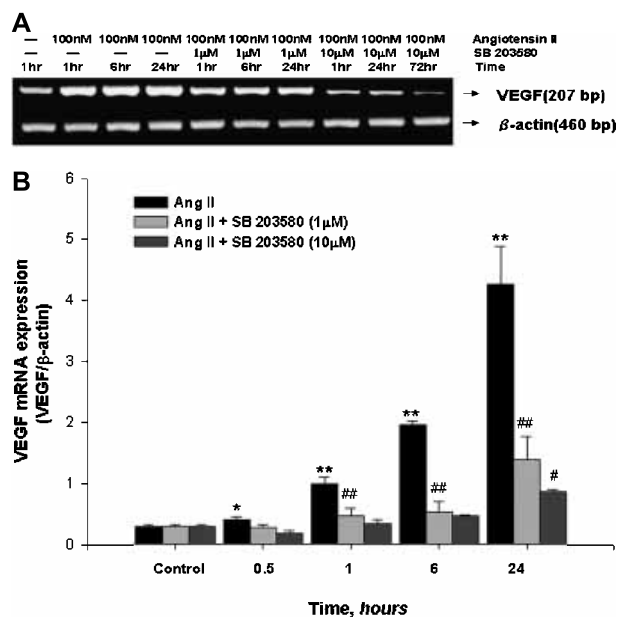


Figure 6 Effects of p38 MAPK inhibitor (SB203580) on the expression of VEGF mRNA induced by Ang-II in cultured podocytes. (A) Representative RT-PCR product. Podocytes were exposed to 100 nM Ang-II with or without different concentrations of SB203580 for 30 min, 1 h, 6 h and 24 h. (B) Densitometric analysis of RT-PCR data. Results are expressed as an optical density ratio of VEGF/ β -actin. Data are shown as means \pm S.D. of 3 independent experiments with triplicate dishes. * P < 0.05, ** P < 0.01 vs control. # P < 0.05 between SB203580-treated groups. ## P < 0.01 vs group treated with 100 nM Ang-II.

Effects of p38 MAPK inhibitor (SB203580) on VEGF protein and mRNA expression increased by Ang-II stimulation

We next examined whether p38 MAPK is involved in Ang-II-induced VEGF production since p38 MAPK, MKK3/6 and CREB were activated by Ang-II treatment. Thus, the inhibitory effect of SB203580 on Ang-II-induced VEGF synthesis was evaluated. With increasing concentrations of SB203580, Ang-II-induced VEGF mRNA expression significantly decreased, indicating that SB203580 acts as an inhibitor of VEGF synthesis. At a concentration of 10 μ M SB203580, VEGF mRNA expression was markedly suppressed from 30 min in response to 100 nM Ang-II (30 min: Ang-II 0.40 ± 0.03 vs Ang-II+SB203580 0.17 ± 0.05 ; 1 h: Ang-II 0.98 ± 0.11 vs Ang-II+SB203580 0.33 ± 0.06 ; 6 h: Ang-II 1.95 ± 0.07 vs Ang-II+SB203580 0.46 ± 0.02 ; 24 h: Ang-II 4.25 ± 0.62 vs Ang-II+SB203580 0.86 ± 0.02) (Fig. 6).

With western blot analysis, VEGF protein synthesis from podocytes also showed significant inhibition by pre-treatment with SB203580 from 30 min. Densitometric analysis demonstrated that SB203580 treatment

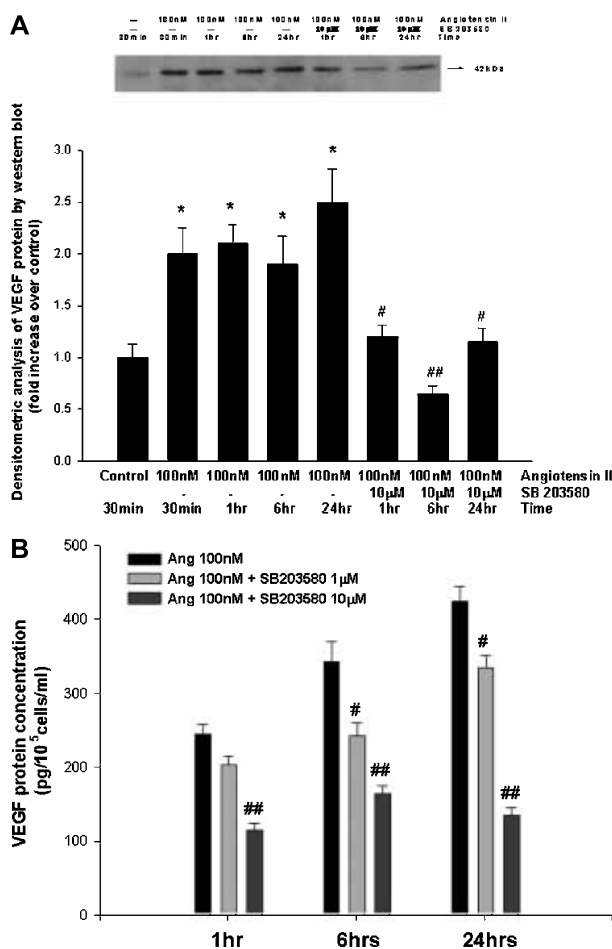


Figure 7 Effects of p38 MAPK inhibitor (SB203580) on the synthesis of VEGF protein in cultured podocytes. Podocytes were exposed to 100 nM Ang-II with or without different concentrations of SB203580 for 30 min, 1 h, 6 h, and 24 h. (A) Representative western blot analysis showing the effect of p38 MAPK inhibitor (SB203580) on the cell content of VEGF protein in cultured podocytes. Thirty micrograms proteins were electrophoresed on 10% sodium dodecyl sulfate-polyacrylamide gel. (B) Secretory VEGF proteins were measured in culture supernatants using ELISA, which recognizes both the 164 and 120 amino acid residue forms of mouse VEGF. Densitometric data are shown as means \pm S.D. of 3 independent experiments with triplicate dishes. * P < 0.05 vs control. # P < 0.05, ## P < 0.01 vs group treated with 100 nM Ang-II.

induced a reduction of 42% at 1 h, 66% at 6 h, and 54% at 24 h in VEGF protein synthesis induced by Ang-II (Fig. 7A).

To confirm the inhibitory effect of SB203580 on VEGF protein release, secretory types of VEGF were measured by the ELISA technique. As illustrated in Fig. 7B, SB203580 pretreatment prevented Ang-II-induced VEGF release in a concentration-dependent manner. Supplementation of culture media with

SB203580 even at a concentration of 1 μ M showed significant inhibition of VEGF release. The levels of VEGF release in the group with prior treatment with SB203580 (10 μ M) were significantly inhibited at 1 h (Ang-II 244 \pm 13 pg/10⁵ cells/ml vs Ang-II+SB203580 114 \pm 9 pg/10⁵ cells/ml), 6 h (Ang-II 342 \pm 28 pg/10⁵ cells/ml vs Ang-II+SB203580 163 \pm 11 pg/10⁵ cells/ml), and 24 h (Ang-II 424 \pm 20 pg/10⁵ cells/ml vs Ang-II+SB203580 134 \pm 11 pg/10⁵ cells/ml).

Effect of Ang-II on electrophoretic mobility shift of CRE elements

To confirm that Ang-II-induced phosphorylation of the p38 MAPK family proteins actually promotes the binding affinity of CREB, gel mobility-shift assays were performed by using an oligonucleotide probe containing the consensus sequence of CRE. When podocytes were treated with 100 nM Ang-II, the CRE DNA-protein complex formation, (the binding of nuclear protein to CRE oligonucleotide containing consensus CRE (5'-TGACGTCA-3')), was increased in a time-dependent manner (Fig. 8A). Ang-II-induced phosphorylation of CREB was consistent with increased binding affinity of CREB (Fig. 8B). Anti-CREB antibody caused supershift, indicating the presence of CREB protein within the CRE DNA-protein complexes (Fig. 8C, lanes 1–4). ³²P-Labeled CRE oligonucleotide was incubated with nuclear extracts in the presence of 0.2-, 2- and 20-fold molar excess of unlabeled competitor. Ang-II-induced protein binding to the ³²P-labeled CRE probe was inhibited by the unlabeled oligonucleotide containing the CRE sequence but not that containing the Oct-1 sequence (Fig. 8C, lanes 5–8). These findings suggest that angiotensin II enhances CRE-directed transcription of VEGF gene through increasing the DNA binding activity and phosphorylation of CREB.

Discussion

Among the many potential pathogenetic mechanisms responsible for the development of diabetic kidney disease, VEGF has been proposed to play a role in the development of diabetic renal changes in type 1 and type 2 diabetes (Duh & Aiello 1999). More direct evidence of VEGF as a potential mediator of diabetic nephropathy was gained from studies using VEGF antibodies. Treatment with VEGF antibody was shown to ameliorate both the classical early features of diabetic renal disease, i.e. renal/glomerular hypertrophy and hyperfiltration and also, more importantly, late renal changes (i.e. basement membrane thickening), with a tendency to reduce total mesangial volume in type 1 and 2 experimental models (De Vriese *et al.* 2001, Flyvbjerg *et al.* 2002).

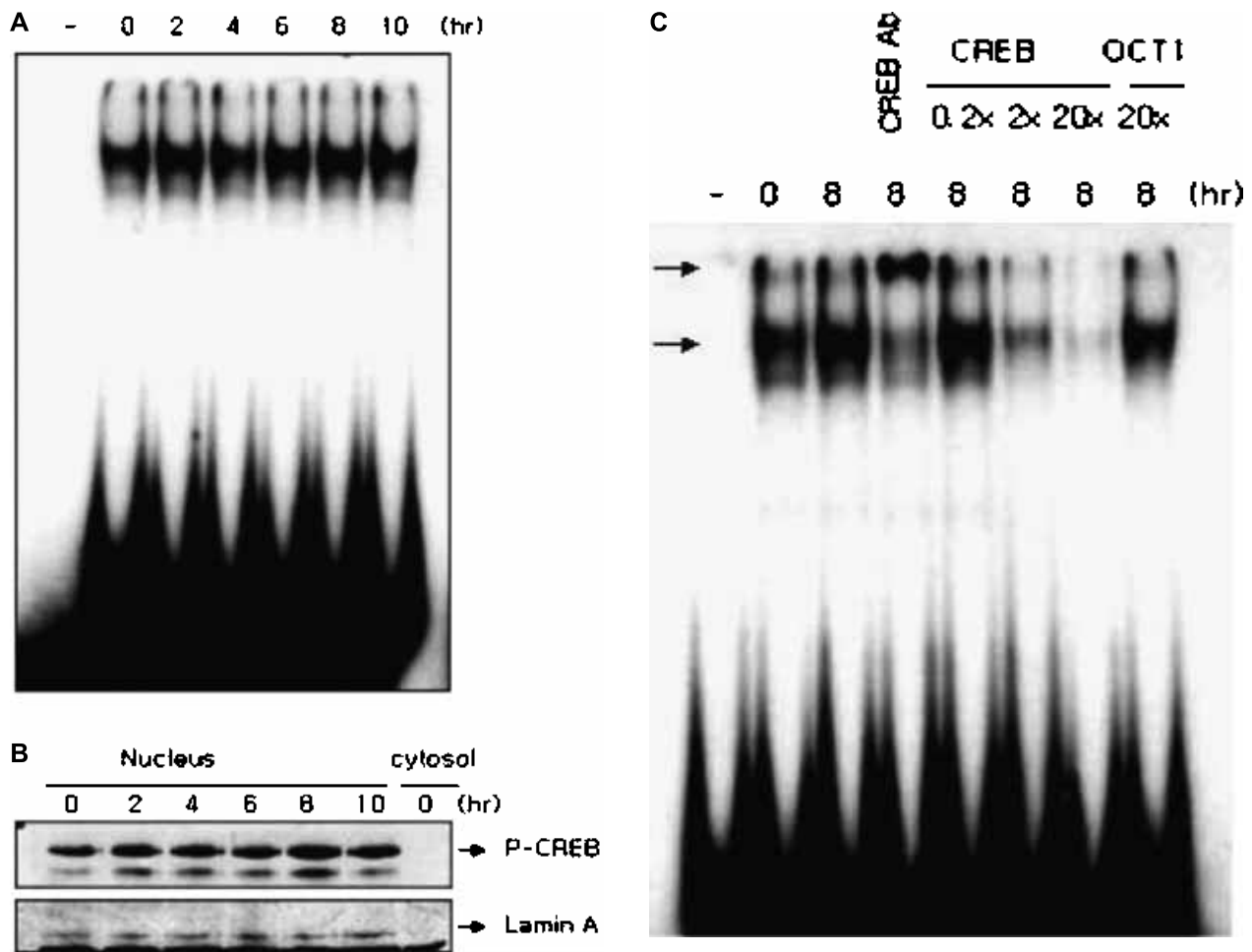


Figure 8 Effect of Ang-II on the binding affinity of CRE-binding protein. (A) Podocytes were treated with 100 nM Ang-II for 2 h, 4 h, 6 h, 8 h and 10 h. Nuclear extract was incubated with 32 P-labeled oligonucleotide containing the consensus cAMP response element (CRE) sequence, and the mixture was separated on a 4% nondenaturing polyacrylamide gel. (B) Podocytes were treated with 100 nM Ang-II for 2 h, 4 h, 6 h, 8 h and 10 h. Nuclear protein was extracted and immunoblot was performed using phospho-specific CREB antibody. (C) Podocytes were treated with 100 nM Ang-II for 8 h. The supershift assay was performed by incubating nuclear extract with or without antibody against CREB. 32 P-Labeled CRE oligonucleotide was incubated with nuclear extracts in the presence of 0.2-, 2- and 20-fold molar excess of unlabeled competitor. The supershifted band is indicated by an arrow (upper band).

We have previously demonstrated that expression of VEGF increased in podocytes in diabetic patients in the early phase of diabetic kidney disease, and that urinary excretion of VEGF significantly increased according to the degree of proteinuria in both human and diabetic rats (Cha *et al.* 2000, 2004).

In the present study, VEGF mRNA expression increased after exposure to Ang-II in a concentration-dependent manner. This finding is similar to previous studies in mesangial cells and vascular smooth muscle cells in response to Ang-II (Williams *et al.* 1995, Kyriakis & Avruch 2001). However, VEGF mRNA expression peaked at 24 h and then gradually decreased.

That Ang-II causes an early, but not sustained, release of VEGF in this study seems to argue against a role for VEGF in diabetic nephropathy, but prolonged activation of renin-angiotensin systems in the diabetic state *in vivo* may induce sustained VEGF overproduction for a longer period.

The effect of Ang-II on VEGF synthesis was maximal at an Ang-II concentration of 100 nM. The concentration of Ang-II required for the maximal effects on VEGF mRNA and protein secretion are consistent with previous reports in vascular smooth muscle cells and mesangial cells (Anderson *et al.* 1993, Williams *et al.* 1995). Similar to the findings with mRNA expression,

VEGF protein content in podocytes and VEGF release also increased in a dose- and time-dependent manner. In addition, we observed that VEGF transcriptional activity also increased after stimulation with Ang-II in a dose-dependent manner. This result is in line with increased VEGF gene expression and protein production after stimulation with Ang-II. Interestingly, the stimulatory effects of Ang-II on VEGF mRNA and protein production were abolished not by pretreatment with PD123319, which is an AT₂ receptor antagonist, but by pretreatment with losartan, which suggests that the AT₁ receptor is involved in Ang-II-mediated VEGF synthesis.

p38 MAPK is a member of the MAPK family and plays an essential role in regulating many cellular processes, including inflammation, cell differentiation, cell growth and death (Ono & Han 2000, Tian *et al.* 2000, Kyriakis & Avruch 2001). There are many reports that Ang-II activates p38 MAPK in various kinds of cells including mesangial cells (Reddy *et al.* 2002) and vascular smooth muscle cells (Touyz *et al.* 2001). There is now a wealth of data supporting a direct role for p38 MAPK as a signaling pathway in the production of VEGF (Tanaka *et al.* 2000, Jung *et al.* 2001, Duyndam *et al.* 2003, Tokuda *et al.* 2003, Tsai *et al.* 2003). Therefore, in this study, we focused on the p38 MAPK pathway to identify the signaling pathways that mediate Ang-II-induced VEGF synthesis.

Since Ang-II stimulated VEGF production in a time-dependent manner with a peak level at the 24-h interval, we examined whether Ang-II activates the p38 MAPK pathway up to 24 h. The activation of p38 MAPK occurred in response to Ang-II after 1 h, peaking at 6 h, and then gradually decreased. Although p38 MAPK activation began to decline at 24 h, it was maintained at a significant level. In addition, Ang-II rapidly induced the activation of MKK3/6 and CREB.

Since the VEGF gene contains a CRE in its promoter region, activation of CREB can induce the transcript for VEGF (Shima *et al.* 1996). Braun *et al.* (2001) reported that the hypoxia responsive element and CRE are equally involved in the regulation of VEGF expression in various tissues in diabetic rats.

In our experiments, activation of CREB also occurred at 30 min of exposure, with the peak occurring at the 24-h interval following incubation. We also demonstrated that Ang-II augmented CREB DNA binding affinity in podocytes associated with rapid phosphorylation of CREB. Taken together, these results suggest that Ang-II activated the p38 MAPK pathway including CREB phosphorylation in podocytes.

Finally, we also examined the possible role of p38 MAPK in Ang-II-induced VEGF production. Pretreatment with the p38 MAPK inhibitor, SB203580, resulted in a dose-dependent decrease in Ang-II-induced VEGF mRNA transcription. SB203580 is widely used as a

potent inhibitor of p38 MAPK and its primary activity is to inhibit p38 MAPK activity by binding to the ATP acceptor pocket (Young *et al.* 1997). However, it has recently been reported that SB203580 can also inhibit the phosphatidylinositol 3-kinase/protein kinase B pathway, cyclin G-associated kinase, and other apoptosis regulatory protein kinases (Lali *et al.* 2000, Godl *et al.* 2003). Thus, our results suggest that Ang-II-induced VEGF expression in podocytes is mediated, in part, through the p38 MAPK pathway. Furthermore, Ang-II-induced VEGF protein production was significantly suppressed by SB203580, suggesting that an Ang-II-p38 MAPK-VEGF pathway may exist and may be active in podocytes.

Although we did not observe whether Ang-II could increase VEGF production by podocytes *in vivo*, several studies have suggested a close relationship between the renin-angiotensin system (RAS) and VEGF in experimental animal models of diabetes. Treatment with Ang-II type 1 receptor antagonist attenuated renal structural injury and suppressed the VEGF expression in the glomeruli in streptozotocin-induced diabetic rats (Qin *et al.* 2003). In addition, angiotensin converting enzyme inhibitor prevented diabetes-induced podocyte injuries such as a decrease in podocyte number and foot process broadening, which results in structural and functional alterations in both type 1 and type 2 diabetic rats (Mifsud *et al.* 2001, Gross *et al.* 2003). Thus, we can postulate that Ang-II is involved in VEGF overexpression which accompanies diabetic nephropathy, and our *in vitro* data provide a potential mechanism for the beneficial effect of RAS blockade in diabetic nephropathy.

In conclusion, the present study provides evidence that Ang-II stimulated VEGF synthesis in cultured mouse podocytes, and that p38 MAPK activity increased after stimulation with Ang-II in association with both MKK3/6 and CREB activation. Furthermore, treatment with a potent p38 MAPK inhibitor abolished Ang-II-mediated VEGF production. Taken together, these results suggest that Ang-II-induced VEGF production was mediated, in part, through the p38 MAPK pathway in cultured mouse podocytes.

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